DAG/TMH:jam 3/3/06 479831 598014D PATENT

Attorney Reference Number 7158-71253-10 Application Number 10/648,631

LISTING OF CLAIMS

This listing of claims will replace all prior versions, and listings of claims in the application:

- 1-3. (Canceled).
- 4. (Currently amended) A substantially pure Pin1 polypeptide comprising consisting of:
- a WW domain consisting of amino acid residues 5-43 of SEQ ID NO:2, or a functional fragment thereof, wherein the WW domain or functional fragment thereof binds to NIMA; and a PPlase domain having peptidyl prolyl isomerase activity.
- 5. (Previously presented) The substantially pure Pin1 polypeptide of claim 4, wherein the peptidyl prolyl isomerase activity of the PPIase domain is not inhibited by cyclosporin A or FK520.
 - 6-7. (Canceled).
- 8. (Currently amended) A substantially pure Pin1 polypeptide comprising consisting of:
- a PPIase domain consisting of amino acid residues 59-163 of SEQ ID NO:2, or a functional fragment thereof, wherein the PPIase domain or functional fragment thereof has peptidyl prolyl isomerase activity; and
 - a WW domain that binds to NIMA.
 - 9-20. (Canceled).
- 21. (Previously presented) A substantially pure WW domain of a Pin1 polypeptide consisting of amino acid residues 5-43 of SEQ ID NO:2, or a functional fragment thereof which binds to NIMA.

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22-24. (Canceled).

25. (Currently amended) The polypeptide of claim 24, wherein the heterologous polypeptide is A substantially pure WW domain of a Pin1 polypeptide consisting of amino acid residues 5-43 of SEQ ID NO:2, or a functional fragment thereof which binds to NIMA, fused to an epitope tag, a carrier protein, a DNA binding domain, a transactivation domain, or an enzyme suitable for use as a label.

26-27. (Canceled).

28. (Previously presented) A substantially pure PPIase domain of a Pin1 polypeptide consisting of amino acid residues 59-163 of SEQ ID NO:2, or a functional fragment thereof having peptidyl prolyl isomerase activity.

29-31. (Canceled).

- 32. (Currently amended) The polypeptide of claim 31, wherein the heterologous polypeptide is A substantially pure PPIase domain of a Pin1 polypeptide consisting of amino acid residues 59-163 of SEO ID NO:2, or a functional fragment thereof having peptidyl prolyl isomerase activity, fused to an epitope tag, a carrier protein, a DNA binding domain, a transactivation domain, or an enzyme suitable for use as a label.
- 33. (Previously presented) A substantially pure fragment of a Pin1 polypeptide, which Pin1 polypeptide consists of amino acid residue 1-163 of SEQ ID NO:2, wherein the fragment comprises amino acid residues 5-43 of SEQ ID NO:2 and binds to NIMA.
- 34. (Previously presented) A substantially pure fragment of a Pin1 polypeptide, which Pin1 polypeptide consists of amino acid residue 1-163 of SEQ ID NO:2, wherein the fragment

NO:2; and

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comprises amino acid residues 59-163 of SEQ ID NO:2 and has peptidyl prolyl isomerase activity.

35. (Currently amended) A substantially pure polypeptide comprising:
a-fragment of a Pin1 polypeptide, which Pin1 polypeptide consists of amino acid residue
1-163 of SEQ ID NO:2, wherein the fragment comprises amino acid 5-43 of SEQ ID NO:2; and
a-heterologous polypeptide,
fused to an epitope tag, a carrier protein, a DNA binding domain, a transactivation domain, or an
enzyme suitable for use as a label.
36. (Currently amended) A substantially pure polypeptide comprising:
a-fragment of a Pin1 polypeptide, which Pin1 polypeptide consists of amino acid residue

wherein the fragment has peptidyl prolyl isomerase activity; and wherein the fragment is fused to an epitope tag, a carrier protein, a DNA binding domain, a transactivation domain, or an enzyme suitable for use as a label.

1-163 of SEQ ID NO:2, wherein the fragment comprises amino acid residues 59-163 of SEQ ID